

ISPA-5001-C1 (Seq List).ST25
SEQUENCE LISTING

<110> Syrrx, Inc.
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Leu Ser Arg Phe Ile Ala Pro Leu Pro Phe Gln Asn Thr Pro Val Val
35 40 45

Glu Thr Met Gln Tyr Gly Ala Leu Leu Gly Gly Lys Arg Leu Arg Pro
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Phe Leu Val Tyr Ala Thr Gly His Met Phe Gly Val Ser Thr Asn Thr
65 70 75 80

Leu Asp Ala Pro Ala Ala Ala Val Glu Cys Ile His Ala Tyr Ser Leu
85 90 95

Ile His Asp Asp Leu Pro Ala Met Asp Asp Asp Asp Leu Arg Arg Gly
100 105 110

Leu Pro Thr Cys His Val Lys Phe Gly Glu Ala Asn Ala Ile Leu Ala
115 120 125

Gly Asp Ala Leu Gln Thr Leu Ala Phe Ser Ile Leu Ser Asp Ala Asp
130 135 140

Met Pro Glu Val Ser Asp Arg Asp Arg Ile Ser Met Ile Ser Glu Leu
145 150 155 160

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Ala Ser Ala Ser Gly Ile Ala Gly Met Cys Gly Gly Gln Ala Leu Asp
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Leu Asp Ala Glu Gly Lys His Val Pro Leu Asp Ala Leu Glu Arg Ile
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His Arg His Lys Thr Gly Ala Leu Ile Arg Ala Ala Val Arg Leu Gly
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Ala Leu Ser Ala Gly Asp Lys Gly Arg Arg Ala Leu Pro Val Leu Asp
210 215 220

Lys Tyr Ala Glu Ser Ile Gly Leu Ala Phe Gln Val Gln Asp Asp Ile
225 230 235 240

Leu Asp Val Val Gly Asp Thr Ala Thr Leu Gly Lys Arg Gln Gly Ala
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Asp Gln Gln Leu Gly Lys Ser Thr Tyr Pro Ala Leu Leu Gly Leu Glu
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Gln Ala Arg Lys Lys Ala Arg Asp Leu Ile Asp Asp Ala Arg Gln Ser
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Leu Ala Asp Tyr Ile Ile Gln Arg Asn Lys
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cgcctgcgac cttccctggt ttatgccacc ggtcatatgt tcggcgtag cacaacacg 240

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| gcgagcgcca gtggtattgc cggaatgtgc ggtggtcagg cattagattt agacgcggaa | 540 |
| ggcaaacacg tacctctgga cgcgctttag cgtattcatc gtcataaaac cggcgcattg | 600 |
| attcgcgccc ccgttcgcct tggtgcatta agcgccggag ataaaggacg tcgtgctcg | 660 |
| ccggtactcg acaagtatgc agagagcatc ggccttgcct tccaggttca ggatgacatc | 720 |
| ctggatgtgg tgggagatac tgcaacgttgg gaaaaacgcc agggtgcga ccagcaactt | 780 |
| ggtaaaagta cctaccctgc acttctgggt cttgagcaag cccggaagaa agcccggat | 840 |
| ctgatcgacg atgcccgtca gtcgctgaaa caactggctg aacagtcaact cgataacctcg | 900 |
| gcacttggaaag cgcttagcgga ctacatcatc cagcgtataa aataa | 945 |